

PCT

RAW SEQUENCE LISTING DATE: 02/11/2003 PATENT APPLICATION: US/10/018,248A TIME: 14:35:12

Input Set : A:\10018248a.raw.txt

Output Set: N:\CRF4\02112003\J018248A.raw

3 <110> APPLICANT: EXELIXIS, INC. 5 <120> TITLE OF INVENTION: ANIMAL MODELS AND METHODS FOR ANALYSIS OF LIPID METABOLISM AND SCREENING OF PHARMACEUTICAL AND PESTICIDAL AGENTS THAT MODULATE LIPID METABOLISM 6 8 <130> FILE REFERENCE: EX99-004C-US 10 <140> CURRENT APPLICATION NUMBER: 10/018,248A C--> 11 <141> CURRENT FILING DATE: 2002-07-30 13 <150> PRIOR APPLICATION NUMBER: US 60/189,700 14 <151> PRIOR FILING DATE: 2000-03-15 16 <150> PRIOR APPLICATION NUMBER: US 09/332,522 17 <151> PRIOR FILING DATE: 1999-06-14 19 <160> NUMBER OF SEQ ID NOS: 8 ENTERED 21 <170> SOFTWARE: PatentIn version 3.1 23 <210> SEO ID NO: 1 24 <211> LENGTH: 3419 25 <212> TYPE: DNA 26 <213> ORGANISM: Caenorhabditis elegans 28 <400> SEQUENCE: 1 29 ggtttaatta cccaagtttg agaatgaacg aagaattcga gggagacgtc cctatgtcgg 60 31 atccgtttct ctcattggtc acaaaattgg atgatattgc gccatttcca aataacgacc 120 33 cgctcgattt tgacatggag cacaactggc aagagcccgg accatcacaa caaccggatc 180 35 catcaattcc cggaaatcaa cacagtccgc cacaggaata ttatgatatt gatggtcaac 240 37 gagacgtaag caccttacac teeetgetea accaeaacaa egaegaette tteteaatge 300 39 gattttcccc gccaaacttt gatctcggcg gaggccgtgg accttctcta gccgccaccc 360 41 aacaattatc tggagaaggt cctgcaagta tgcttaaccc cttacaaaca tctccaccaa 420 43 gtggaggtta ccccccggca gatgcctaca gacctctatc acttqctcaa caactcqccq 480 45 cgccagcgat gactccacat caggcagcgt cgctttttgt taatactaat ggaattgatc 540 600 49 catatacaga agccatggga catatcaacg ggtacatgtc tccatacgac caagctcaag 660 51 gcccatcagg accatcatat tactcacaac accatcaatc tccaccacct catcaccacc 720 53 atcaccaccc gatgccaaaa atccatgaga accctgaaca agtggcatct ccatcgattg 780 55 aagatgetee agagaegaaa eeaacteatt tggttgaace acaaagteea aaaageeege 840 57 agaatatgaa agaggagctt cttcggttac tagttaacat gtctccgagt gaagttgaac 900 59 ggttaaagaa taaaaaatca ggagcatgtt cagcgacgaa tgggccatcg aggagtaagg 960 61 agaaggcggc gaagattgtg attcaggaga cagcggaagg ggatgaagat gaggatgatg 1020 63 aggatagtga ttccggggag actatgtctc agggaactac tattattgtt cgaagaccaa 1080 65 aaaccgagcg tcgtacggca cacaatctca tcgaaaagaa gtatagatgc tcaataaatg 1140 67 atcgaattca acagctgaaa gtacttttgt gtggggatga agctaagctt tcaaaatcgg 1200 69 caacactacg acgggctatt gaacatatcg aggaggttga acacgagaat caggtgttga 1260 71 agcatcatgt tgaacaaatg agaaagacac tgcagaataa tcgattaccg tacccggaac 1320

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Input Set : A:\10018248a.raw.txt

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173 Glu Tyr Tyr Asp Ile Asp Gly Gln Arg Asp Val Ser Thr Leu His Ser
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Input Set : A:\10018248a.raw.txt

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	ьeu	Leu	ASII	птѕ		Asn	ASP	ASP	rne		ser	мес	Arg	Pne		PLO
178	D	7\	Dha	71	85	C1	C1	C1	7\ ~~ ~~	90	D	C	т	7.1.	95	mb ~
	Pro	ASI	Pne	_	ьeu	Gly	СТУ	СТУ	_	СТУ	Pro	ser	ьeu		Ата	1111
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	GIn	GIn		Ser	GTA	Glu	GTA		Ата	Ser	мет	ьeu		Pro	ьeu	GIN
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	Thr		Pro	Pro	Ser	Gly	_	Tyr	Pro	Pro	Ala		Ата	Tyr	Arg	Pro
190	_	130	_			- •	135			_		140		_		
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Input Set : A:\10018248a.raw.txt

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	Ser	Pro		GIU	His	GTÀ	Arg		тте	Asp	Asp	Pro	_	GIA	Thr	Ser
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		TT ! -	77-7	T	m1		<b>01</b>	T 3 -	m	m		<b>.</b>	70	<b>.</b>	<b>.</b>			
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W>	422 <b>425</b> 426 429 430 433 434 437	Thr Val Phe	1055 Trp 1070 Arg 1085 His 1100 >> SE	Ser Gly Leu	Gly Arg His	Xaa Val Thr	Arg Ser	106 Arc 107 Met 109 Val	50 75 A1 90	i <b>l A</b> i	r <b>g S</b> e	<b>er T</b> h	1( nr Ly 1( la Gl 1( nr Se	065 <b>/s h</b> 080 ln H 095	<b>Met</b> 2	<b>Asp</b> .	<b>Ala</b> Ala	
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W>	422 425 426 429 430 433 434 437 438 439 440 442 443 445	Thr Val Phe <210 <211 <212 <213 <400 cggc tcgg	1055 Trp 1070 Arg 1085 His 1100 > SE > TY 3 > OR cacga ggacc	Ser  Gly  Leu  QID  NGTH  PE:  GANI  QUEN  gg a  ag a	Gly Arg His NO: 19 DNA SM: CE: ttaa ggag	Val Thr 3 71 Dros 3 tgct taag	Ser Leu ophi g at	106 Arc 107 Met 109 Val 110 la m	of Variation Var	nl Andreg Andres Lee	eg Se	er Ther Alis There	10 11 12 12 13 14 17 17 17 17 17 17 17 17 17 17 17 17 17	065 ys 1 080 In H 095 er 1 110	Met A	Asp Asp Asp gtat	Ala Ala Leu	120 180
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W>	422 425 426 429 430 433 434 437 438 439 440 442 443 445 447 449 451	Thr Val Phe <210 <211 <212 <213 <400 cgga tcgc tcgc tagt	1055 Trp 1070 Arg 1085 His 1100 > SE > TY 3> OR caggaco aggaco agtg caactg	Gly Leu QID NGTH PE: GANI QUEN gg a ag a ag a ag a	Arg His NO: 19 DNA SM: CE: ttaa ggag tcaa tcgc	Val Thr 3 71 Dros 3 tgct taag gcaa gaca ttta	Ser Lew ophi g att c cgt t gg	106 Arg 107 Met 109 Val 110 la m ttct tcgg	y Var 55 Ar 60 Ly 05 Ar 190 Ly 195 Ar 190 Ar	al Ar	eg Seeu Historia	er Ther Alis There acac egte gtag ecat ecga	agca cggt aatc	065  ys N 080  ln H 095  er N 110  ettgg caggg cccct	Met A	Asp Asp Asp dtattactactactactagtgt	Ala Ala Leu aaggag agcatt tcctaa tcttca gttgca	120 180 240 300
W>	422 425 426 429 430 433 434 437 438 439 440 442 443 445 447 449 451 453	Thr Val Phe <210 <211 <212 <213 <400 cgga tcga tcaattagt	1055 Trp 1070 Arg 1085 His 1100 > SE > TY 3 > OR cacga gacca actga cactga cactga	Gly Leu QID NGTH PE: GANI QUEN gg a ag a ag a ag a ag a	Arg His NO: 19 DNA SM: CE: ttaa ggag tcaa tcgc tacg	Val Thr 3 71 Dros 3 tgct taag gcaa gcaa gttta atgc	Ser Lew ophi g att ctt gg	106 Arg 107 Met 109 Val 110  la m ttct tcgg gagg tgag cgtt cct c	y Var 55 An 60 Ly 55 Ly 60 Ly	al Ar	eg Seeu Historia	er There Alis The acac egte grag ecat ecga greet	agca cggt aatc	065  ys 1 080  In I 095  er 1 110  ettgo	Met A	Asp Asp Asp datatactactactactactactactactactactact	Ala Ala Leu aaggag agcatt tcctaa tcttca gttgca tgagcc	120 180 240 300 360
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	422 425 426 429 430 433 434 437 438 449 440 445 447 449 451 453 457 461	Thr Val Phe <210<<211<<212<<400 cgga tcga tcca tgca tgct tgct gct	1055 Trp 1070 Arg 1085 His 1100 > SE > TY 3> OR cacga cactg caccg cactg caccg	Ser  Gly  Leu  QID  NGTH  PE:  GANI  QUEN  gg a  ac g  cc g  cc g  cc g  ct t	Arg His NO: 19 DNA SM: ttaa ggag tcaat tcgc acga acga aggtc ggtat	Val Thr 3 71 Dros tgaaa gaaa ttta ggtaat ttgat tgat	Ser Leu ophi attccgcttttttt	106 Arg 107 Met 109 Val 110 la m ttcggagg tgagg	y Va y Va y S y S y S y S y S y S y S y S	al Ar	g Se seu History ster sacta actory ster sacta actory steet at get at get at get at get at get actory sectory s	er There Allis The acac gtag gctag gacg gacg gacg gacg gac	agca cggt agca ccggt aatc gagt ccgg	of the second of	Met A	Asp Asp gtattactagtgaagattcagaacct	Ala Ala Leu aaggag agcatt tcctaa tcttca gttgca tgagcc gtaaca tttctc aagaca ttccc atgccg	120 180 240 300 360 420 480 540
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Input Set : A:\10018248a.raw.txt

Output Set: N:\CRF4\02112003\J018248A.raw

#### Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 1073
Seq#:7; N Pos. 3038,3049
Seq#:8; Xaa Pos. 993

#### Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5